File Copy
09/466778
STN

```
=> d his

     (FILE 'HOME' ENTERED AT 13:15:51 ON 15 APR 2001)

     FILE 'BIOSIS, CAPLUS, EMBASE, MEDLINE, SCISEARCH' ENTERED AT 13:16:36 ON
     15 APR 2001
L1             3 S FELL PROTEIN
L2           388 S HYALURONATE (P)(BINDING (W) PROTEIN)
L3           584 S L2 OR HABP
L4             0 S L3 (P) FELL
L5             0 S L3 AND L1
L6            30 S L3 (P) CD44
L7             0 S L3 (P) ((CD44 (S) PRECURSOR))
L8             0 S L3 (P) ((CD44 (W)LIKE))

=> log off y
```

09/466778

	Type	L #	Hits	Search Text	DBs	Time Stamp	Comments	Error Definition	Errors
1	BRS	L1	0	FELL adj protein	USPAT; US-PGPUB; EPO; JPO; DERWENT	2001/04/15 12:31			0
2	BRS	L7	23	CD44 same precursor	USPAT; US-PGPUB; EPO; JPO; DERWENT	2001/04/15 12:38			0
3	BRS	L19	2	L13 same CD44	USPAT; US-PGPUB; EPO; JPO; DERWENT	2001/04/15 12:41			0
4	BRS	L13	16	hyaluronate with (binding adj protein)	USPAT; US-PGPUB; EPO; JPO; DERWENT	2001/04/15 12:50			0

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EAST Search Strategy

	Type	L #	Hits	Search Text	DBs	Time Stamp	Comments	Error Definition	Errors
5	BRS	L25	26	L13 or HABP	USPAT; US-PGPUB; EPO; JPO; DERWENT	2001/04/15 12:51			0

(85)

NCBI

National Center for Biotechnology Information

National Library of Medicine National Institutes of Health

PubMed Entrez BLAST OMIM Taxonomy Structure

Search GenBank for Go

SITE MAP

About NCBI
general and contact information

GenBank
sequence submission support and software

Molecular databases
sequences, structures and taxonomy

Literature databases
PubMed and OMIM

Genomic biology
the human genome, whole genomes and related resources

Tools
for data mining

Research at NCBI
people, projects and seminars

Education

What does NCBI do?

Established in 1988 as a national resource for molecular biology information, NCBI creates public databases, conducts research in computational biology, develops software tools for analyzing genome data, and disseminates biomedical information - all for the better understanding of molecular processes affecting human health and disease.

Draft Human Genome

Explore human genome resources or browse the human genome sequence using the Map Viewer.

Integrated STS reports

Archives

UniSTS

UniSTS presents marker information collected from public resources including GenBank, RHdb, GDB, and various maps. Zero in on primer and mapping data, e-PCR results, and Map Viewer and LocusLink cross-references. More...

NCBI in the News

The draft sequences of the human genome were compared in an article by Aach et al. (*Nature*, Feb.

Hot Spots

- Cancer genome anatomy project
- Clusters of orthologous groups
- Coffee Break
- Electronic PCR
- Gene expression omnibus
- Genes and disease
- Human genome resources
- Human/mouse homology maps
- LocusLink
- Malaria genetics & genomics
- ORF finder
- Reference sequence project

teaching resources and on-line tutorials

FTP site
download data and software

15), noting overall similarities but differences in details. Use of RefSeq, NCBI's manually curated database of mRNA sequences, is credited with providing additional gene annotation for the public sequence.

Revised April 4, 2001

Retrovirus resources

- Serial analysis of gene expression
- Trace archive
- UniGene
- VecScreen

Credits

Mirror sites

Jobs

About GeneCards
Disease genes
How to Mirror
Guiding the user
Data Sources
Data Extraction
Usage statistics
Citing this resource
Publications

What's New

Version: 2.19
Release: Jan 31, 2001
Entries: 18,583
Approved*:11,980

About Bioinformatics
Data Mining in Biology
Web Usability
Science on the Web

Your Feedback

GeneCards: human genes, maps, proteins and diseases

GeneCards is a database of human genes, their products and their involvement in diseases. It offers **concise information** about the **functions** of all **human genes** that have an approved symbol, as well as selected others [gene listing].

GeneCards now also supports searching UDB (The Unified Database for Human Genome Mapping). *Read more about UDB.*

Search

[Quick Start] [Guided Tour] [More search examples]

- **Search/Display GeneCards by** keyword(s)

  Go

  For example, you can display the GeneCard for the (case-sensitive) **symbol** BRCA1,
  or search GeneCards for the **keyword(s)**:
  - p53
  - apolipoprot***AND** (hyper* **OR** Alzheim*)
  - U85267 *GenBank accession No.*
  - Hs.1288 *UniGene cluster*
  - ATCC:106253, image:303124 *clone identifier*
  - chromosome: 22, locus: 20p*,locus: 7p13

- **Search UDB integrated map**
  specify chromosome: 1 Go
  For example, a map region in chromosome 19.

- **Search UDB by gene/marker name** Go
  For example, Information about the mapped marker D17S1843.

- **View estimated boundaries (in Megabases) of cytogenetic bands**
  specify chromosome: 1 Go

What's special about GeneCards?

The information presented here has been **automatically extracted** from various resources by scripts developed in our group. GeneCards is

particulary useful for people who wish to find information about genes of interest in the context of functional genomics and proteomics.

This resource also features a new type of **navigation support** system that **guides** its users to the information. Important parts of this guidance system are the **spell corrector**, and the automatically generated tips for query reformulation.

Developed at the **Crown Human Genome Center** & **Bioinformatics Unit**, at the **Weizmann Institute of Science**

Credits:

Michael Rebhan, Avital Adato, Vered Chalifa-Caspi, Inga Peter, Jaime Prilusky,
Michal Ronen, Hershel Safer, Marilyn Safran, Shai Shen-Orr , Liora Yaar,
Doron Lancet

Comments to: *cards@bioinfo.weizmann.ac.il*

* *Entries with HUGO-approved symbols.*

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